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<110> Quip Technology Limited

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<141> 2000-08-16

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<151> 2000-08-16

<160> 19

<170> PatentIn version 3.1

<210> 1

<211> 524

<212> PRT

<213> porcine endogenous retrovirus

<400> 1

Met Gly Gln Thr Val Thr Thr Pro Leu Ser Leu Thr Leu Asp His Trp 10 15

Thr Glu Val Arg Ser Arg Ala His Asn Leu Ser Val Gln Val Lys Lys 20 25 30

Gly Pro Trp Gln Thr Phe Cys Ala Ser Glu Trp Pro Thr Phe Asp Val

Gly Trp Pro Ser Glu Gly Thr Phe Asn Ser Glu Ile Ile Leu Ala Val 50 60

Lys Ala Ile Ile Phe Gln Thr Gly Pro Gly Ser His Pro Asp Gln Glu 65 70 75 80 Pro Tyr Ile Leu Thr Trp Gln Asp Leu Ala Glu Asp Pro Pro Pro Trp 85 90 95 Val Lys Pro Trp Leu Asn Lys Pro Arg Lys Pro Gly Pro Arg Ile Leu 100 105 110 Ala Leu Gly Glu Lys Asn Lys His Ser Ala Glu Lys Val Glu Pro Ser 115 120 125 Ser Ser Tyr Leu Pro Arg Asp Arg Gly Ala Ala Asp Leu Ala Gly Thr 130 140 Pro Thr Cys Ser Pro Thr Pro Leu Ser Ser Thr Gly Cys Cys Glu Gly 145 150 155 160 Thr Ser Ala Pro Pro Gly Ala Pro Val Val Glu Gly Pro Ala Ala Gly
165 170 175 Thr Arg Ser Arg Arg Gly Ala Thr Pro Glu Arg Thr Asp Glu Ile Ala 180 185 190 Ile Leu Pro Leu Arg Thr Tyr Gly Pro Pro Met Pro Gly Gly Gln Leu 195 200 205 Gln Pro Leu Gln Tyr Trp Pro Phe Ser Ser Ala Asp Leu Tyr Asn Trp 210 225 Lys Thr Asn His Pro Pro Phe Ser Glu Asp Pro Gln Arg Leu Thr Gly 225 230 235 240 Leu Val Glu Ser Leu Met Phe Ser His Gln Pro Thr Trp Asp Asp Cys 245 250 255 Gln Gln Leu Leu Gln Thr Leu Phe Thr Thr Glu Glu Arg Glu Arg Ile 260 265 270 Leu Leu Glu Ala Arg Lys Asn Val Pro Gly Ala Asp Gly Arg Pro Thr 275 280 285 Gln Leu Gln Asn Glu Ile Asp Met Gly Phe Pro Leu Thr Arg Pro Gly 290 295 300 Trp Asp Tyr Asn Thr Ala Glu Gly Arg Glu Ser Leu Lys Ile Tyr Arg 305 310 315 320 Gln Ala Leu Val Ala Gly Leu Arg Gly Ala Ser Arg Arg Pro Thr Asn 325 330 335 Leu Ala Lys Val Arg Glu Val Met Gln Gly Pro Asn Glu Pro Pro Ser 340 350

Val Phe Leu Glu Arg Leu Met Glu Ala Phe Arg Arg Phe Thr Pro Phe 355 360 365

Asp Pro Thr Ser Glu Ala Gln Lys Ala Ser Val Ala Leu Ala Phe Ile 370 375 380

Gly Gln Ser Ala Leu Asp Ile Arg Lys Lys Leu Gln Arg Leu Glu Gly 385 390 395

Leu Gln Glu Ala Glu Leu Arg Asp Leu Val Arg Glu Ala Glu Lys Val 405 410 415

Tyr Tyr Arg Arg Glu Thr Glu Glu Glu Lys Glu Gln Arg Lys Glu Lys 420 425 430

Glu Arg Glu Glu Arg Glu Glu Arg Arg Arg Arg Gln Glu Lys Asn 435 440 445

Leu Thr Lys Ile Leu Ala Ala Val Val Glu Gly Lys Ser Ser Arg Glu 450 460

Arg Glu Arg Asp Phe Arg Lys Ile Arg Ser Gly Pro Arg Gln Ser Gly 465 470 475 480

Asn Leu Gly Asn Arg Thr Pro Leu Asp Lys Asp Gln Cys Ala Tyr Cys 485 490 495

Lys Glu Lys Gly His Trp Ala Arg Asn Cys Pro Lys Lys Gly Asn Lys 500 510

Gly Pro Lys Val Leu Ala Leu Glu Glu Asp Lys Asp 515 520

<210> 2

<211> 660

<212≤ PPT

<213> porcine endogenous retrovirus Type PERV A

<400> 2

Met His Pro Thr Leu Ser Arg Arg His Leu Pro Ile Arg Gly Gly Lys
5 10 15

Pro Lys Arg Leu Lys Ile Pro Leu Ser Phe Ala Ser Ile Ala Trp Phe 20 25 30

Leu Thr Leu Ser Ile Thr Pro Gln Val Asn Gly Lys Arg Leu Val Asp 35 40 45

Ser Pro Asn Ser His Lys Pro Leu Ser Leu Thr Trp Leu Leu Thr Asp 50 55 60 Ser Gly Thr Gly Ile Asn Ile Asn Ser Thr Gln Gly Glu Ala Pro Leu 65 70 75 80 Gly Thr Trp Trp Pro Glu Leu Tyr Val Cys Leu Arg Ser Val Ile Pro 85 90 95 Gly Leu Asn Asp Gln Ala Thr Pro Pro Asp Val Leu Arg Ala Tyr Gly
100 105 110 Phe Tyr Val Cys Pro Gly Pro Pro Asn Asn Glu Glu Tyr Cys Gly Asn 115 120 125 Pro Gln Asp Phe Phe Cys Lys Gln Trp Ser Cys Ile Thr Ser Asn Asp 130 135 140 Gly Asn Trp Lys Trp Pro Val Ser Gln Gln Asp Arg Val Ser Tyr Ser 145 150 155 Phe Val Asn Asn Pro Thr Ser Tyr Asn Gln Phe Asn Tyr Gly His Gly 165 170 Arg Trp Lys Asp Trp Gln Gln Arg Val Gln Lys Asp Val Arg Asn Lys
180
185 Gln Ile Ser Cys His Ser Leu Asp Leu Asp Tyr Leu Lys Ile Ser Phe 195 200 Thr Glu Lys Gly Lys Gln Glu Asn Ile Gln Lys Trp Val Asn Gly Ile 210 215 220 Ser Trp Gly Ile Val Tyr Tyr Gly Gly Ser Gly Arg Lys Lys Gly Ser 225 230 235 Val Leu Thr Ile Arg Leu Arg Ile Glu Thr Gln Met Glu Pro Pro Val 245 250 255 Ala Ile Gly Pro Asn Lys Gly Leu Ala Glu Gln Gly Pro Pro Ile Gln 260 270 Glu Gln Arg Pro Ser Pro Asn Pro Ser Asp Tyr Asn Thr Thr Ser Gly 275 280 285 Ser Val Pro Thr Glu Pro Asn Ile Thr Ile Lys Thr Gly Ala Lys Leu 290 295 300 Phe Ser Leu Ile Gln Gly Ala Phe Gln Ala Leu Asn Ser Thr Thr Pro 305 310 315 320 Glu Ala Thr Ser Ser Cys Trp Leu Cys Leu Ala Ser Gly Pro Pro Tyr 325 330 335

Tyr Glu Gly Met Ala Arg Gly Gly Lys Phe Asn Val Thr Lys Glu His 340 345 350

Arg Asp Gln Cys Thr Trp Gly Ser Gln Asn Lys Leu Thr Leu Thr Glu 355 360 365

Val Ser Gly Lys Gly Thr Cys Ile Gly Met Val Pro Pro Ser His Gln 370 380

His Leu Cys Asn His Thr Glu Ala Phe Asn Arg Thr Ser Glu Ser Gln 385 390 395 400

Tyr Leu Val Pro Gly Tyr Asp Arg Trp Trp Ala Cys Asn Thr Gly Leu 405 410 415

Thr Pro Cys Val Ser Thr Leu Val Phe Asn Gln Thr Lys Asp Phe Cys 420 425 430

Val Met Val Gln Ile Val Pro Arg Val Tyr Tyr Pro Glu Lys Ala 435 440 445

Val Leu Asp Glu Tyr Asp Tyr Arg Tyr Asn Arg Pro Lys Arg Glu Pro 450 460

Ile Ser Leu Thr Leu Ala Val Met Leu Gly Leu Gly Val Ala Ala Gly 465 470 475 480

Val Gly Thr Gly Thr Ala Ala Leu Ile Thr Gly Pro Gln Gln Leu Glu 485 490 495

Lys Gly Leu Ser Asn Leu His Arg Ile Val Thr Glu Asp Leu Gln Ala 500 510

Leu Glu Lys Ser Val Ser Asn Leu Glu Glu Ser Leu Thr Ser Leu Ser 515 520 525

Glu Val Val Leu Gln Asn Arg Arg Gly Leu Asp Leu Leu Phe Leu Lys 530 540

Glu Gly Gly Leu Cys Val Ala Leu Lys Glu Glu Cys Cys Phe Tyr Val 545 550 555 560

Asp His Ser Gly Ala Ile Arg Asp Ser Met Ser Lys Leu Arg Glu Arg 565 570

Leu Glu Arg Arg Arg Glu Arg Glu Ala Asp Gln Gly Trp Phe Glu 580 585 590

Gly Trp Phe Asn Arg Ser Pro Trp Met Thr Thr Leu Leu Ser Ala Leu 595 600 605

Thr Gly Pro Leu Val Val Leu Leu Leu Leu Leu Thr Val Gly Pro Cys 610 620

Leu Ile Asn Arg Phe Val Ala Phe Val Arg Glu Arg Val Ser Ala Val 625 635 640

Gln Ile Met Val Leu Arg Gln Gln Tyr Gln Gly Leu Leu Ser Gln Gly 645 650 655

Glu Thr, Asp Leu

<210> 3

<211> 638

<212> PRT

<213> porcine endogenous retrovirus Type PERV MSL

<400> 3

Met His Pro Thr Leu Asn Arg Arg His Leu Pro Ile Arg Gly Gly Lys
1 10 15

Pro Lys Arg Leu Lys Ile Pro Leu Ser Phe Ala Ser Ile Ala Trp Phe 20 25 30

Leu Thr Leu Ser Ile Thr Ser Gln Thr Asn Gly Met Arg Ile Gly Asp 35 40 45

Ser Leu Asn Ser His Lys Pro Leu Ser Leu Thr Trp Leu Ile Thr Asp 50 60

Ser Gly Thr Gly Ile Asn Ile Asn Asn Thr Gln Gly Glu Ala Pro Leu 65 70 75 80

Gly Thr Trp Trp Pro Asp Leu Tyr Val Cys Leu Arg Ser Val Ile Pro 85 90 95

Ser Leu Thr Ser Pro Pro Asp Ile Leu His Ala His Gly Phe Tyr Val $100 \hspace{1cm} 105 \hspace{1cm} 110$

Cys Pro Gly Pro Pro Asn Asn Gly Lys His Cys Gly Asn Pro Arg Asp 115 120 125

Phe Phe Cys Lys Gln Trp Asn Cys Val Thr Ser Asn Asp Gly Tyr Trp 130 140

Lys Trp Pro Thr Ser Gln Gln Asp Arg Val Ser Phe Ser Tyr Val Asn 145 150 155 160

Thr Tyr Thr Ser Ser Gly Gln Phe Asn Tyr Leu Thr Trp Ile Arg Thr 165 170 175

Gly Ser Pro Lys Cys Ser Pro Ser Asp Leu Asp Tyr Leu Lys Ile Ser 180 185 190

Phe Thr Glu Lys Gly Lys Gln Glu Asn Ile Leu Lys Trp Val Asn Gly 195 200 205

Met Ser Trp Gly Met Val Tyr Tyr Gly Gly Ser Gly Lys Gln Pro Gly 210 215 220

Ser Ile Leu Thr Ile Arg Leu Lys Ile Asn Gln Leu Glu Pro Pro Met 225 230 235 240

Ala Ile Gly Pro Asn Thr Val Leu Thr Gly Gln Arg Pro Pro Thr Gln 245 250 255

Gly Pro Gly Pro Ser Ser Asn Ile Thr Ser Gly Ser Asp Pro Thr Glu 260 265 270

Ser Asn Ser Thr Thr Lys Met Gly Ala Lys Leu Phe Ser Leu Ile Gln 275 280 285

Gly Ala Phe Gln Ala Leu Asn Ser Thr Thr Pro Glu Ala Thr Ser Ser 290 295 300

Cys Trp Leu Cys Leu Ala Ser Gly Pro Pro Tyr Tyr Glu Gly Met Ala 305 310 315 320

Arg Arg Gly Lys Phe Asn Val Thr Lys Glu His Arg Asp Gln Cys Thr 325 330 335

Trp Gly Ser Gln Asn Lys Leu Thr Leu Thr Glu Val Ser Gly Lys Gly 340 345 350

Thr Cys Ile Gly Lys Val Pro Pro Ser His Gln His Leu Cys Asn His 355 360 365

Thr Glu Ala Phe Asn Gln Thr Ser Glu Ser Gln Tyr Leu Val Pro Gly 370 380

Tyr Asp Arg Trp Trp Ala Cys Asn Thr Gly Leu Thr Pro Cys Val Ser 385 390 395 400

Thr Leu Val Phe Asn Gln Thr Lys Asp Phe Cys Ile Met Val Gln Ile 405 410 415

Val Pro Arg Val Tyr Tyr Pro Glu Lys Ala Ile Leu Asp Glu Tyr 420 425 430

Asp Tyr Arg Asn His Arg Gln Lys Arg Glu Pro Ile Ser Leu Thr Leu 435 440 445 Ala Val Met Leu Gly Leu Gly Val Ala Ala Gly Val Gly Thr Gly Thr 450 460

Ala Ala Leu Val Thr Gly Pro Gln Gln Leu Glu Thr Gly Leu Ser Asn 465 470 475 480

Leu His Arg Ile Val Thr Glu Asp Leu Gln Ala Leu Glu Lys Ser Val 485 490 495

Ser Asn Leu Glu Glu Ser Leu Thr Ser Leu Ser Glu Val Val Leu Gln
500 510

Asn Arg Arg Gly Leu Asp Leu Leu Phe Leu Lys Glu Gly Gly Leu Cys 515 520

Val Ala Leu Lys Glu Glu Cys Cys Phe Tyr Val Asp His Ser Gly Ala 530 540

Ile Arg Asp Ser Met Asn Lys Leu Arg Glu Arg Leu Glu Lys Arg Arg 545 550 555 560

Arg Glu Lys Glu Thr Thr Gln Gly Trp Phe Glu Gly Trp Phe Asn Arg 565 570 575

Ser Leu Trp Leu Ala Thr Leu Leu Ser Ala Leu Thr Gly Pro Leu Ile 580 585 590

Val Leu Leu Leu Leu Thr Val Gly Pro Cys Ile Ile Asn Lys Leu 595 600 605

Ile Ala Phe Ile Arg Glu Arg Ile Ser Ala Val Gln Ile Met Val Leu 610 615 620

Arg Gln Gln Tyr Gln Ser Pro Ser Ser Arg Glu Ala Gly Arg 625 630 635

<210> 4

<211> 638

<212> PRT

<213> porcine endogenous retrovirus Type Tsukuba

<400> 4

Met His Pro Thr Leu Ser Arg Arg His Leu Pro Ile Arg Gly Gly Lys
10 15

Pro Lys Arg Leu Lys Ile Pro Leu Ser Phe Ala Ser Ile Ala Trp Phe 20 25 30

Leu Thr Leu Ser Ile Thr Ser Gln Thr Asn Gly Met Arg Ile Gly Asp 35 40 45

Ser	Leu 50	Asn	Ser	His	Lys	Pro 55	Leu	Ser	Leu	Thr	Trp 60	Leu	Ile	Thr	Asp

- Ser Gly Thr Gly Ile Asn Ile Asn Asn Thr Gln Gly Glu Ala Pro Leu 65 70 75 80
- Gly Thr Trp Trp Pro Asp Leu Tyr Val Cys Leu Arg Ser Val Ile Pro 85 90 95
- Ser Leu Thr Ser Pro Pro Asp Ile Leu His Ala His Gly Phe Tyr Val 100 105 110
- Cys Pro Gly Pro Pro Asn Asn Gly Lys His Cys Gly Asn Pro Arg Asp 115 120 125
- Phe Phe Cys Lys Gln Trp Asn Cys Val Thr Ser Asn Asp Gly Tyr Trp 130 140
- Lys Trp Pro Thr Ser Gln Gln Asp Arg Val Ser Phe Ser Tyr Val Asn 150 155 160
- Thr Tyr Thr Ser Ser Gly Gln Phe Asn Tyr Leu Thr Trp Ile Arg Thr 165 170 175
- Gly Ser Pro Lys Cys Ser Pro Ser Asp Leu Asp Tyr Leu Lys Ile Ser 180 185 190
- Phe Thr Glu Lys Gly Lys Gln Glu Asn Ile Leu Lys Trp Val Asn Gly 195 205
- Met Ser Trp Gly Met Val Tyr Tyr Gly Gly Ser Gly Lys Gln Pro Gly 210 220
- Ser Ile Leu Thr Ile Arg Leu Lys Ile Asn Gln Leu Glu Pro Pro Met 235 235
- Ala Ile Gly Pro Asn Thr Val Leu Thr Gly Gln Arg Pro Pro Thr Gln 245 250
- Gly Pro Gly Pro Ser Ser Asn Ile Thr Ser Gly Ser Asp Pro Thr Glu 260 265 270
- Ser Ser Ser Thr Thr Lys Met Gly Ala Lys Leu Phe Ser Leu Ile Gln 285
- Gly Ala Phe Gln Ala Leu Asn Ser Thr Thr Pro Glu Ala Thr Ser Ser 290 300
- Cys Trp Leu Cys Leu Ala Leu Gly Pro Pro Tyr Tyr Glu Gly Met Ala 310 315

Arg Arg Gly Lys Phe Asn Val Thr Lys Glu His Arg Asp Gln Cys Thr 325 330 335

Trp Gly Ser Gln Asn Lys Leu Thr Leu Thr Glu Val Ser Gly Lys Gly 340 345

Thr Cys Ile Gly Lys Val Pro Pro Ser His Gln His Leu Cys Asn His 355 360 365

Thr Glu Ala Phe Asn Gln Thr Ser Glu Ser Gln Tyr Leu Val Pro Gly 370 380

Tyr Asp Arg Trp Trp Ala Cys Asn Thr Gly Leu Thr Pro Cys Val Ser 385 390 395 400

Thr Leu Val Phe Asn Gln Thr Lys Asp Phe Cys Ile Met Val Gln Ile 405 410 415

Val Pro Arg Val Tyr Tyr Pro Glu Lys Ala Ile Leu Asp Glu Tyr 420 425 430

Asp Tyr Arg Asn His Arg Gln Lys Arg Glu Pro Ile Ser Leu Thr Leu 435 440 445

Ala Val Met Leu Gly Leu Gly Val Ala Ala Gly Val Gly Thr Gly Thr 450 460

Ala Ala Leu Val Thr Gly Pro Gln Gln Leu Glu Thr Gly Leu Ser Asn 465 470 475 480

Leu His Arg Ile Val Thr Glu Asp Leu Gln Ala Leu Glu Lys Ser Val 485 490 495

Ser Asn Leu Glu Glu Ser Leu Thr Ser Leu Ser Glu Val Val Leu Gln
500 510

Asn Arg Arg Gly Leu Asp Leu Leu Phe Leu Lys Glu Gly Gly Leu Cys 515 520 525

Val Ala Leu Lys Glu Glu Cys Cys Phe Tyr Val Asp His Ser Gly Ala 530 540

Ile Arg Asp Ser Met Asn Lys Leu Arg Glu Arg Leu Glu Lys Arg Arg 545 550 555 560

Arg Glu Lys Glu Thr Thr Gln Gly Trp Phe Glu Gly Trp Phe Asn Arg 565 570 575

Ser Pro Trp Leu Ala Thr Leu Leu Ser Ala Leu Thr Gly Pro Leu Ile 580 585 590

Val Leu Leu Leu Leu Thr Val Gly Pro Cys Ile Ile Asn Lys Leu 595 600 605

Ile Ala Phe Ile Arg Glu Arg Ile Ser Ala Val Gln Ile Met Val Leu 610 620

Arg Gln Gln Tyr Gln Ser Pro Ser Ser Arg Glu Ala Gly Arg 625 630 635

<210> 5

<211> 657

<212> PRT

<213> porcine endogenous retrovirus Type PERV B

<400> 5

Met His Pro Thr Leu Ser Trp Arg His Leu Pro Thr Arg Gly Gly Glu 10 15

Pro Lys Arg Leu Arg Ile Pro Leu Ser Phe Ala Ser Ile Ala Trp Phe 20 25 30

Leu Thr Leu Thr Ile Thr Pro Gln Ala Ser Ser Lys Arg Leu Ile Asp 40 45

Ser Ser Asn Pro His Arg Pro Leu Ser Leu Thr Trp Leu Ile Ile Asp 50 60

Pro Asp Thr Gly Val Thr Val Asn Ser Thr Arg Gly Val Ala Pro Arg 65 75 80

Gly Thr Trp Trp Pro Glu Leu His Phe Cys Leu Arg Leu Ile Asn Pro 85 90 95

Ala Val Lys Ser Thr Pro Pro Asn Leu Val Arg Ser Tyr Gly Phe Tyr 100 105 110

Cys Cys Pro Gly Thr Glu Lys Glu Lys Tyr Cys Gly Gly Ser Gly Glu 115 120 125

Ser Phe Cys Arg Arg Trp Ser Cys Val Thr Ser Asn Asp Gly Asp Trp 130 135

Lys Trp Pro Ile Ser Leu Gln Asp Arg Val Lys Phe Ser Phe Val Asn 150 155 160

Ser Gly Pro Gly Lys Tyr Lys Val Met Lys Leu Tyr Lys Asp Lys Ser 165 170 175

Cys Ser Pro Ser Asp Leu Asp Tyr Leu Lys Ile Ser Phe Thr Glu Lys 180 185 190 Gly Lys Gln Glu Asn Ile Gln Lys Trp Ile Asn Gly Met Ser Trp Gly
195 200 205

Ile Val Phe Tyr Lys Tyr Gly Gly Gly Ala Gly Ser Thr Leu Thr Ile 210 215 220

Arg Leu Arg Ile Glu Thr Gly Thr Glu Pro Pro Val Ala Val Gly Pro 235 230 235

Asp Lys Val Leu Ala Glu Gln Gly Pro Pro Ala Leu Glu Pro Pro His 245 250 255

Asn Leu Pro Val Pro Gln Leu Thr Ser Leu Arg Pro Asp Ile Thr Gln 265 270

Pro Pro Ser Asn Gly Thr Thr Gly Leu Ile Pro Thr Asn Thr Pro Arg 275 280 285

Asn Ser Pro Gly Val Pro Val Lys Thr Gly Gln Arg Leu Phe Ser Leu 290 300

Ile Gln Gly Ala Phe Gln Ala Ile Asn Ser Thr Asp Pro Asp Ala Thr 305 310 315

Ser Ser Cys Trp Leu Cys Leu Ser Ser Gly Pro Pro Tyr Tyr Glu Gly 325 335

Met Ala Lys Glu Gly Lys Phe Asn Val Thr Lys Glu His Arg Asn Gln 340 350

Cys Thr Trp Gly Ser Arg Asn Lys Leu Thr Leu Thr Glu Val Ser Gly 365

Lys Gly Thr Cys Ile Gly Lys Ala Pro Pro Ser His Gln His Leu Cys 375 380

Tyr Ser Thr Val Val Tyr Glu Gln Ala Ser Glu Asn Gln Tyr Leu Val 385 390 395

Pro Gly Tyr Asn Arg Trp Trp Ala Cys Asn Thr Gly Leu Thr Pro Cys 410 415

Val Ser Thr Ser Val Phe Asn Gln Ser Lys Asp Phe Cys Val Met Val 420 425 430

Gln Ile Val Pro Arg Val Tyr Tyr His Pro Glu Glu Val Val Leu Asp 435 440 445

Glu Tyr Asp Tyr Arg Tyr Asn Arg Pro Lys Arg Glu Pro Val Ser Leu 450 460

Thr Leu Ala Val Met Leu Gly Leu Gly Thr Ala Val Gly Val Gly Thr 465 470 475 480 Gly Thr Ala Ala Leu Ile Thr Gly Pro Gln Gln Leu Glu Lys Gly Leu 485 490 495

Gly Glu Leu His Ala Ala Met Thr Glu Asp Leu Arg Ala Leu Glu Glu 500 510

Ser Val Ser Asn Leu Glu Glu Ser Leu Thr Ser Leu Ser Glu Val Val 515 520

Leu Gln Asn Arg Arg Gly Leu Asp Leu Leu Phe Leu Arg Glu Gly Gly 530

Leu Cys Ala Ala Leu Lys Glu Glu Cys Cys Phe Tyr Val Asp His Ser 550 555

Gly Ala Ile Arg Asp Ser Met Ser Lys Leu Arg Glu Arg Leu Glu Arg 565 570 575

Arg Arg Arg Glu Arg Glu Ala Asp Gln Gly Trp Phe Glu Gly Trp Phe 580 590

Asn Arg Ser Pro Trp Met Thr Thr Leu Leu Ser Ala Leu Thr Gly Pro 605

Leu Val Val Leu Leu Leu Leu Thr Val Gly Pro Cys Leu Ile Asn 610 615

Arg Phe Val Ala Phe Val Arg Glu Arg Val Ser Ala Val Gln Ile Met 625 635 640

Val Leu Arg Gln Gln Tyr Gln Gly Leu Leu Ser Gln Gly Glu Thr Asp 650 655

Leu

<210> 6

<211> 656

<212> PRT

<213> porcine endogenous retrovirus Type PoEV1

<400> 6

Met His Pro Thr Leu Ser Arg Arg His Leu Pro Thr Arg Gly Glu 10 15

Pro Lys Arg Leu Arg Ile Pro Leu Ser Phe Ala Ser Ile Ala Trp Phe 20 30

Leu Thr Leu Thr Ile Thr Pro Gln Ala Ser Ser Lys Arg Leu Ile Asp 35 40 45

Ser Ser Asn Pro His Arg Pro Leu Ser Leu Thr Trp Leu Ile Ile Asp 50 55 60

Pro Asp Thr Gly Val Thr Val Asn Ser Thr Arg Gly Val Ala Pro Arg 65 70 75 80

Gly Thr Trp Trp Pro Glu Leu His Phe Cys Leu Arg Leu Ile Asn Pro 85 90 95

Ala Val Lys Ser Thr Pro Pro Asn Leu Val Arg Ser Tyr Gly Phe Tyr 100 105 110

Cys Cys Pro Gly Thr Glu Lys Glu Lys Tyr Cys Gly Gly Ser Gly Glu 115 120 125

Ser Phe Cys Arg Arg Trp Ser Cys Val Thr Ser Asn Asp Gly Asp Trp 130 140

Lys Trp Pro Ile Ser Leu Gln Asp Arg Val Lys Phe Ser Phe Val Asn 145 150 155 160

Ser Gly Pro Gly Lys Tyr Lys Met Met Lys Leu Tyr Lys Asp Lys Ser 165 170 175

Cys Ser Pro Ser Asp Leu Asp Tyr Leu Lys Ile Ser Phe Thr Glu Arg 180 185 190

Lys Thr Gly Lys Tyr Ser Lys Val Asp Lys Trp Tyr Glu Leu Gly Asn 195 200 205

Ser Phe Leu Leu Tyr Gly Gly Gly Ala Gly Ser Thr Leu Thr Ile Arg 210 215 220

Leu Arg Ile Glu Thr Gly Thr Glu Pro Pro Val Ala Met Gly Pro Asp 225 230 235

Lys Val Leu Ala Glu Gln Gly Pro Pro Ala Leu Glu Pro Pro His Asn 245 250 255

Leu Pro Val Pro Gln Leu Thr Ser Leu Arg Pro Asp Ile Thr Gln Pro 260 265 270

Pro Ser Asn Ser Thr Thr Gly Leu Ile Pro Thr Asn Thr Pro Arg Asn 275 280 285

Ser Pro Gly Val Pro Val Lys Thr Gly Gln Arg Leu Phe Ser Leu Ile 290 295 300

Gln Gly Ala Phe Gln Ala Ile Asn Ser Thr Asp Pro Asp Ala Thr Ser 305 310 315 320

Ser Cys Trp Leu Cys Leu Ser Ser Gly Pro Pro Tyr Tyr Glu Gly Met 325 330 335 Ala Lys Glu Arg Lys Phe Asn Val Thr Lys Glu His Arg Asn Gln Cys 340 345 350 Thr Trp Gly Ser Arg Asn Lys Leu Thr Leu Thr Glu Val Ser Gly Lys 355 360 365 Gly Thr Cys Ile Gly Lys Ala Pro Pro Ser His Gln His Leu Cys Tyr 370 380 Ser Thr Val Val Tyr Glu Gln Ala Ser Glu Asn Gln Tyr Leu Val Pro 385 390 395 400 Gly Tyr Asn Arg Trp Trp Ala Cys Asn Thr Gly Leu Thr Pro Cys Val 405 410 415 Ser Thr Ser Val Phe Asn Gln Ser Lys Asp Phe Cys Val Met Val Gln 420 430 Ile Val Pro Arg Val Tyr Tyr His Pro Glu Glu Val Val Leu Asp Glu
435 440 445 Tyr Asp Tyr Arg Tyr Asn Arg Pro Lys Arg Glu Pro Val Ser Leu Thr 450 460 Leu Ala Val Met Leu Gly Leu Gly Thr Ala Val Gly Val Gly Thr Gly 465 470 475 480 Thr Ala Ala Leu Ile Thr Gly Pro Gln Gln Leu Glu Lys Gly Leu Gly 495 Glu Leu His Ala Ala Met Thr Glu Asp Leu Arg Ala Leu Lys Glu Ser 500 510 Val Ser Asn Leu Glu Glu Ser Leu Thr Ser Leu Ser Glu Val Val Leu 515 520 525 Gln Asn Arg Arg Gly Leu Asp Leu Leu Phe Leu Arg Glu Gly Gly Leu 530 540 Cys Ala Ala Leu Lys Glu Glu Cys Cys Phe Tyr Val Asp His Ser Gly 545 550 555 Ala Ile Arg Asp Ser Met Asn Lys Leu Arg Lys Lys Leu Glu Arg Arg 565 570 575 Arg Arg Glu Arg Glu Ala Asp Gln Gly Trp Phe Glu Gly Trp Phe Asn 580 590

Arg Ser Pro Trp Met Thr Thr Leu Leu Ser Ala Leu Thr Gly Pro Leu 595 605

Val Val Leu Leu Leu Leu Thr Val Gly Pro Cys Leu Ile Asn Arg 610 620

Phe Val Ala Phe Val Arg Glu Arg Val Ser Ala Val Gln Ile Met Val 625 630 635

Leu Arg Gln Gln Tyr Gln Gly Leu Leu Ser Gln Gly Glu Thr Asp Leu 645 650 655

<210> 7

<211> 221

<212> PRT

<213> porcine endogenous retrovirus Type PERV A

<400> 7

Asn Asn Glu Glu Tyr Cys Gly Asn Pro Gln Asp Phe Phe Cys Lys Gln 10 15

Trp Ser Cys Ile Thr Ser Asn Asp Gly Asn Trp Lys Trp Pro Val Ser 25 30

Gln Gln Asp Arg Val Ser Tyr Ser Phe Val Asn Asn Pro Thr Ser Tyr 35 45

Asn Gln Phe Asn Tyr Gly His Gly Arg Trp Lys Asp Trp Gln Gln Arg 50 60

Val Gln Lys Asp Val Arg Asn Lys Gln Ile Ser Cys His Ser Leu Asp 70 75 80

Leu Asp Tyr Leu Lys Ile Ser Phe Thr Glu Lys Gly Lys Gln Glu Asn 90 95

Ile Gln Lys Trp Val Asn Gly Ile Ser Trp Gly Ile Val Tyr Tyr Gly
100 105 110

Gly Ser Gly Arg Lys Lys Gly Ser Val Leu Thr Ile Arg Leu Arg Ile 115 120 125

Glu Thr Gln Met Glu Pro Pro Val Ala Ile Gly Pro Asn Lys Gly Leu 130 140

Ala Glu Gln Gly Pro Pro Ile Gln Glu Gln Arg Pro Ser Pro Asn Pro 150 155 160

Ser Asp Tyr Asn Thr Thr Ser Gly Ser Val Pro Thr Glu Pro Asn Ile 165 170 175 Thr Ile Lys Thr Gly Ala Lys Leu Phe Ser Leu Ile Gln Gly Ala Phe 180 185 190

Gln Ala Leu Asn Ser Thr Thr Pro Glu Ala Thr Ser Ser Cys Trp Leu 195 200 205

Cys Leu Ala Ser Gly Pro Pro Tyr Tyr Glu Gly Met Ala 210 215 220

<210> 8

<211> - 203

<212> PRT

<213> porcine endogenous retrovirus Type PERV MSL

<400> .8

Asn Asn Gly Lys His Cys Gly Asn Pro Arg Asp Phe Phe Cys Lys Gln 10 15

Trp Asn Cys Val Thr Ser Asn Asp Gly Tyr Trp Lys Trp Pro Thr Ser 20 25 30

Gln Gln Asp Arg Val Ser Phe Ser Tyr Val Asn Thr Tyr Thr Ser Ser 35 40 45

Gly Gln Phe Asn Tyr Leu Thr Trp Ile Arg Thr Gly Ser Pro Lys Cys 50 60

Ser Pro Ser Asp Leu Asp Tyr Leu Lys Ile Ser Phe Thr Glu Lys Gly 65 70 75 80

Lys Gln Glu Asn Ile Leu Lys Trp Val Asn Gly Met Ser Trp Gly Met 85 90 95

Val Tyr Tyr Gly Gly Ser Gly Lys Gln Pro Gly Ser Ile Leu Thr Ile 100 105 110

Arg Leu Lys Ile Asn Gln Leu Glu Pro Pro Met Ala Ile Gly Pro Asn 115 120 125

Thr Val Leu Thr Gly Gln Arg Pro Pro Thr Gln Gly Pro Gly Pro Ser 130 140

Ser Asn Ile Thr Ser Gly Ser Asp Pro Thr Glu Ser Asn Ser Thr Thr 145 150 155 160

Lys Met Gly Ala Lys Leu Phe Ser Leu Ile Gln Gly Ala Phe Gln Ala 165 170 175 Leu Asn Ser Thr Thr Pro Glu Ala Thr Ser Ser Cys Trp Leu Cys Leu 180 185 190

Ala Ser Gly Pro Pro Tyr Tyr Glu Gly Met Ala 195 200

<210> 9

<211> 203

<212> PRT

<213> porcine endogenous retróvirus Type Tsukuba

<400> 9

Asn Asn Gly Lys His Cys Gly Asn Pro Arg Asp Phe Phe Cys Lys Gln 10 15

Trp Asn Cys Val Thr Ser Asn Asp Gly Tyr Trp Lys Trp Pro Thr Ser 20 25 30

Gln Gln Asp Arg Val Ser Phe Ser Tyr Val Asn Thr Tyr Thr Ser Ser 35 40 45

Gly Gln Phe Asn Tyr Leu Thr Trp Ile Arg Thr Gly Ser Pro Lys Cys 50 60

Ser Pro Ser Asp Leu Asp Tyr Leu Lys Ile Ser Phe Thr Glu Lys Gly 65 70 75 80

Lys Gln Glu Asn Ile Leu Lys Trp Val Asn Gly Met Ser Trp Gly Met 85 90 95

Val Tyr Tyr Gly Gly Ser Gly Lys Gln Pro Gly Ser Ile Leu Thr Ile 100 105 110

Arg Leu Lys Ile Asn Gln Leu Glu Pro Pro Met Ala Ile Gly Pro Asn 115 120 125

Thr Val Leu Thr Gly Gln Arg Pro Pro Thr Gln Gly Pro Gly Pro Ser 130 140

Ser Asn Ile Thr Ser Gly Ser Asp Pro Thr Glu Ser Ser Ser Thr Thr 145 150 155 160

Lys Met Gly Ala Lys Leu Phe Ser Leu Ile Gln Gly Ala Phe Gln Ala 165 170 175

Leu Asn Ser Thr Thr Pro Glu Ala Thr Ser Ser Cys Trp Leu Cys Leu 180 185 190

Ala Leu Gly Pro Pro Tyr Tyr Glu Gly Met Ala 195 200 <210> 10

<211> 221

<212> PRT

<213> porcine endogenous retrovirus Type PERV B

<400> 10

Glu Lys Glu Lys Tyr Cys Gly Gly Ser Gly Glu Ser Phe Cys Arg Arg 1 5 10 15

Trp Ser Cys Val Thr Ser Asn Asp Gly Asp Trp Lys Trp Pro Ile Ser 20 25 30

Leu Gln Asp Arg Val Lys Phe Ser Phe Val Asn Ser Gly Pro Gly Lys 35 40 45

Tyr Lys Val Met Lys Leu Tyr Lys Asp Lys Ser Cys Ser Pro Ser Asp 50 55 60

Leu Asp Tyr Leu Lys Ile Ser Phe Thr Glu Lys Gly Lys Gln Glu Asn 65 70 75 80

Ile Gln Lys Trp Ile Asn Gly Met Ser Trp Gly Ile Val Phe Tyr Lys 85 90 95

Tyr Gly Gly Ala Gly Ser Thr Leu Thr Ile Arg Leu Arg Ile Glu 100 105 110

Thr Gly Thr Glu Pro Pro Val Ala Val Gly Pro Asp Lys Val Leu Ala 115 120 125

Glu Gln Gly Pro Pro Ala Leu Glu Pro Pro His Asn Leu Pro Val Pro 130 140

Gln Leu Thr Ser Leu Arg Pro Asp Ile Thr Gln Pro Pro Ser Asn Gly 155 160

Thr Thr Gly Leu Ile Pro Thr Asn Thr Pro Arg Asn Ser Pro Gly Val 165 170 175

Pro Val Lys Thr Gly Gln Arg Leu Phe Ser Leu Ile Gln Gly Ala Phe 180 185 190

Gln Ala Ile Asn Ser Thr Asp Pro Asp Ala Thr Ser Ser Cys Trp Leu 195 200 205

Cys Leu Ser Ser Gly Pro Pro Tyr Tyr Glu Gly Met Ala 210 215 220 <210> 11

<211> 220

<212> PRT

<213> porcine endogenous retrovirus Type PoEV1

<400> 11

Glu Lys Glu Lys Tyr Cys Gly Gly Ser Gly Glu Ser Phe Cys Arg Arg 1 10 15

Trp Ser Cys Val Thr Ser Asn Asp Gly Asp Trp Lys Trp Pro Ile Ser 20 25 30

Leu Gln Asp Arg Val Lys Phe Ser Phe Val Asn Ser Gly Pro Gly Lys 35 40 45

Tyr Lys Met Met Lys Leu Tyr Lys Asp Lys Ser Cys Ser Pro Ser Asp 50 60

Leu Asp Tyr Leu Lys Ile Ser Phe Thr Glu Arg Lys Thr Gly Lys Tyr 65 75 80

Ser Lys Val Asp Lys Trp Tyr Glu Leu Gly Asn Ser Phe Leu Leu Tyr 85 90 95

Gly Gly Gly Ala Gly Ser Thr Leu Thr Ile Arg Leu Arg Ile Glu Thr $100 \hspace{1cm} 105 \hspace{1cm} 110$

Gly Thr Glu Pro Pro Val Ala Met Gly Pro Asp Lys Val Leu Ala Glu 115 120 125

Gln Gly Pro Pro Ala Leu Glu Pro Pro His Asn Leu Pro Val Pro Gln 130 135 140

Leu Thr Ser Leu Arg Pro Asp Ile Thr Gln Pro Pro Ser Asn Ser Thr 145 150 155 160

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